

=> d his full

(FILE 'HOME' ENTERED AT 13:53:33 ON 12 OCT 2006)

INDEX 'ADISCTI, ADISINSIGHT, ADISNEWS, AGRICOLA, ANABSTR, ANTE, AQUALINE, AQUASCI, BIOENG, BIOSIS, BIOTECHABS, BIOTECHDS, BIOTECHNO, CABA, CAPLUS, CEABA-VTB, CIN, CONFSCI, CROPB, CROPU, DDFB, DDFU, DGENE, DISSABS, DRUGB, DRUGMONOG2, DRUGU, EMBAL, EMBASE, ...' ENTERED AT 13:54:08 ON 12 OCT 2006
SEA DDL? OR (ALANI?(S)LIGAS?)

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185 FILE WPINDEX
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L1 QUE DDL? OR (ALANI?(S) LIGAS?)

D RANK

FILE 'USPATFULL, GENBANK, SCISEARCH, CAPLUS, PROMT, BIOSIS, PASCAL,
EMBASE, MEDLINE, LIFESCI' ENTERED AT 13:56:02 ON 12 OCT 2006

L2 9522 SEA DDL? OR (ALANI?(S) LIGAS?)
L3 501 SEA L2 AND (TUBERCULOS? OR SMEGMAT?)
L4 399 SEA L3 AND (VECTOR? OR PLASMID?)
L5 58 SEA L4 AND GPM?
L6 58 DUP REM L5 (0 DUPLICATES REMOVED)
D TI L6 1-58
L7 172 SEA L1(S)(TUBERCULOS? OR SMEGMAT?)
L8 83 SEA L7 (S) (VECTOR? OR PLASMID?)
L9 17 SEA L8(S)(GPM?)
L10 17 DUP REM L9 (0 DUPLICATES REMOVED)
D TI L10 1-17
L11 38 SEA L8 AND (GPM? OR PBUN?)
L12 38 DUP REM L11 (0 DUPLICATES REMOVED)
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* * * * * Welcome to STN International * * * * *

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NEWS 5 MAY 11 KOREAPAT updates resume
NEWS 6 MAY 19 Derwent World Patents Index to be reloaded and enhanced
NEWS 7 MAY 30 IPC 8 Rolled-up Core codes added to CA/CAPLUS and
USPATFULL/USPAT2
NEWS 8 MAY 30 The F-Term thesaurus is now available in CA/CAPLUS
NEWS 9 JUN 02 The first reclassification of IPC codes now complete in
INPADOC
NEWS 10 JUN 26 TULSA/TULSA2 reloaded and enhanced with new search and
and display fields
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NEWS 14 JUL 19 Coverage of Research Disclosure reinstated in DWPI
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NEWS 17 AUG 30 CA(SM)/CAPLUS(SM) Austrian patent law changes
NEWS 18 SEP 11 CA/CAPLUS enhanced with more pre-1907 records
NEWS 19 SEP 21 CA/CAPLUS fields enhanced with simultaneous left and right
truncation
NEWS 20 SEP 25 CA(SM)/CAPLUS(SM) display of CA Lexicon enhanced
NEWS 21 SEP 25 CAS REGISTRY(SM) no longer includes Concord 3D coordinates
NEWS 22 SEP 25 CAS REGISTRY(SM) updated with amino acid codes for pyrrolysine
NEWS 23 SEP 28 CEABA-VTB classification code fields reloaded with new
classification scheme

NEWS EXPRESS JUNE 30 CURRENT WINDOWS VERSION IS V8.01b, CURRENT
MACINTOSH VERSION IS V6.0c(ENG) AND V6.0Jc(JP),
AND CURRENT DISCOVER FILE IS DATED 26 JUNE 2006.

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FILE 'HOME' ENTERED AT 13:53:33 ON 12 OCT 2006

=> index bioscience medicine

FILE 'DRUGMONOG' ACCESS NOT AUTHORIZED

COST IN U.S. DOLLARS

SINCE FILE	TOTAL
ENTRY	SESSION
0.21	0.21

FULL ESTIMATED COST

INDEX 'ADISCTI, ADISINSIGHT, ADISNEWS, AGRICOLA, ANABSTR, ANTE, AQUALINE, AQUASCI, BIOENG, BIOSIS, BIOTECHABS, BIOTECHDS, BIOTECHNO, CABA, CAPLUS, CEABA-VTB, CIN, CONFSCI, CROPB, CROPU, DDFB, DDFU, DGENE, DISSABS, DRUGB, DRUGMONOG2, DRUGU, EMBAL, EMBASE, ...' ENTERED AT 13:54:08 ON 12 OCT 2006

71 FILES IN THE FILE LIST IN STNINDEX

Enter SET DETAIL ON to see search term postings or to view search error messages that display as 0* with SET DETAIL OFF.

=> s ddl4 or (alani4(s)ligas4)

4) IS NOT A RECOGNIZED COMMAND

The previous command name entered was not recognized by the system.
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=> s ddl? or (alani?(s)ligas?)

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57 FILES HAVE ONE OR MORE ANSWERS, 71 FILES SEARCHED IN STNINDEX

L1 QUE DDL? OR (ALANI?(S) LIGAS?)

=> d rank

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F39	10	RDISCLOSURE
F40	7	ANTE

F41	7	PHIN
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F43	6	DRUGB
F44	6	FROSTI
F45	5	HEALSAFE
F46	5	WATER
F47	3	AQUASCI
F48	3	IMSPRODUCT
F49	3	PHAR
F50	2	AQUALINE
F51	2	IMSRESEARCH
F52	1	ADISINSIGHT
F53	1	DRUGMONOG2
F54	1	PHARMAML
F55	1	PROUSDDR
F56	1	VETU
F57	1	WPIFV

=> file f1-f7, f9-f11

COST IN U.S. DOLLARS

SINCE FILE

ENTRY

TOTAL

SESSION

FULL ESTIMATED COST

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2.04

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=> s ddl? or (alani?(s)ligas?)

L2 9522 DDL? OR (ALANI?(S) LIGAS?)

=> s l2 and (tuberculos? or smegmat?)

L3 501 L2 AND (TUBERCULOS? OR SMEGMAT?)

=> s l3 and (vector? or plasmid?)

L4 399 L3 AND (VECTOR? OR PLASMID?)

=> s l4 and gpm?

L5 58 L4 AND GPM?

=> dup rem l5

DUPLICATE IS NOT AVAILABLE IN 'GENBANK'.

ANSWERS FROM THESE FILES WILL BE CONSIDERED UNIQUE

PROCESSING COMPLETED FOR L5

L6 58 DUP REM L5 (0 DUPLICATES REMOVED)

=> d ti l6 1-58

L6 ANSWER 1 OF 58 USPATFULL on STN

TI Novel polynucleotides

L6 ANSWER 2 OF 58 USPATFULL on STN

TI Hollow fiber technique for in vivo study of cell populations

L6 ANSWER 3 OF 58 USPATFULL on STN

TI Microorganisms for therapy

L6 ANSWER 4 OF 58 USPATFULL on STN

TI Nucleic acid and amino acid sequences relating to Staphylococcus epidermidis for diagnostics and therapeutics

L6 ANSWER 5 OF 58 USPATFULL on STN

TI Nucleic acid and amino acid sequences relating to Enterobacter cloacae for diagnostics and therapeutics

L6 ANSWER 6 OF 58 USPATFULL on STN

TI Methods and materials relating to novel polypeptides and polynucleotides

L6 ANSWER 7 OF 58 USPATFULL on STN

TI Nucleic acid and amino acid sequences relating to streptococcus pneumoniae for diagnostics and therapeutics

L6 ANSWER 8 OF 58 USPATFULL on STN

TI Microorganisms for therapy

L6 ANSWER 9 OF 58 USPATFULL on STN

TI Recombinant mycobacteria overexpressing D-alanine ligase gene and uses therefore

L6 ANSWER 10 OF 58 USPATFULL on STN

TI Therapeutic agents useful for treating pain

L6 ANSWER 11 OF 58 USPATFULL on STN

TI Therapeutic agents useful for treating pain

L6 ANSWER 12 OF 58 USPATFULL on STN

TI Novel nucleic acids and polypeptides

L6 ANSWER 13 OF 58 USPATFULL on STN

TI Novel nucleic acids and polypeptides

L6 ANSWER 14 OF 58 USPATFULL on STN

TI Streptococcus pneumoniae polynucleotides and sequences

L6 ANSWER 15 OF 58 USPATFULL on STN

TI Methods of diagnosis of breast cancer, compositions and methods of screening for modulators of breast cancer

L6 ANSWER 16 OF 58 USPATFULL on STN

TI Novel human polynucleotides and polypeptides encoded thereby

L6 ANSWER 17 OF 58 USPATFULL on STN
 TI Nucleic acid and amino acid sequences relating to Streptococcus pneumoniae for diagnostics and therapeutics

L6 ANSWER 18 OF 58 USPATFULL on STN
 TI Nucleic acid sequences relating to Candida albicans for diagnostics and therapeutics

L6 ANSWER 19 OF 58 USPATFULL on STN
 TI Nucleic acid and amino acid sequences relating to Enterococcus faecalis for diagnostics and therapeutics

L6 ANSWER 20 OF 58 USPATFULL on STN
 TI Nucleic acid sequences and expression system relating to Enterococcus faecium for diagnostics and therapeutics

L6 ANSWER 21 OF 58 USPATFULL on STN
 TI Nucleic acid and amino acid sequences relating to Acinetobacter baumannii for diagnostics and therapeutics

L6 ANSWER 22 OF 58 MEDLINE on STN
 TI Roles of Mycobacterium smegmatis D-alanine:D-alanine ligase and D-alanine racemase in the mechanisms of action of and resistance to the peptidoglycan inhibitor D-cycloserine.

L6 ANSWER 23 OF 58 USPATFULL on STN
 TI Novel Polynucleotides

L6 ANSWER 24 OF 58 USPATFULL on STN
 TI ENTEROCOCCUS FAECALIS POLYNUCLEOTIDES AND POLYPEPTIDES

L6 ANSWER 25 OF 58 USPATFULL on STN
 TI ENTEROCOCCUS FAECALIS POLYNUCLEOTIDES AND POLYPEPTIDES

L6 ANSWER 26 OF 58 USPATFULL on STN
 TI STREPTOCOCCUS PNEUMONIAE POLYNUCLEOTIDES AND SEQUENCES

L6 ANSWER 27 OF 58 GENBANK® COPYRIGHT 2006 on STN
 TITLE (TI): Comparison of the genome sequence of the poultry pathogen Bordetella avium with those of B. bronchiseptica, B. pertussis, and B. parapertussis reveals extensive diversity in surface structures associated with host interaction
 TITLE (TI): Direct Submission

L6 ANSWER 28 OF 58 GENBANK® COPYRIGHT 2006 on STN
 TITLE (TI): The genome of Rhizobium leguminosarum has recognizable core and accessory components
 TITLE (TI): Direct Submission

L6 ANSWER 29 OF 58 GENBANK® COPYRIGHT 2006 on STN
 TITLE (TI): The complete genome sequence of the European Francisella tularensis subspecies tularensis isolate FSC 198 suggests that it is derived from the archetypal laboratory strain Schu S4, originally isolated in North America
 TITLE (TI): Direct Submission

L6 ANSWER 30 OF 58 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Complete sequence of chromosome of Mycobacterium sp.
MCS

TITLE (TI): Direct Submission

L6 ANSWER 31 OF 58 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): The partitioned Rhizobium etli genome: Genetic and
metabolic redundancy in seven interacting replicons

TITLE (TI): Direct Submission

L6 ANSWER 32 OF 58 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): The partitioned Rhizobium etli genome: Genetic and
metabolic redundancy in seven interacting replicons

TITLE (TI): Direct Submission

L6 ANSWER 33 OF 58 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Complete Sequence of Chromosome 1 of Rhodobacter
sphaeroides 2.4.1

TITLE (TI): Direct Submission

L6 ANSWER 34 OF 58 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): The Chlamydophila abortus genome sequence reveals an
array of variable proteins that contribute to
interspecies variation

TITLE (TI): The Chlamydophila abortus genome sequence reveals an
array of variable proteins that contribute to
interspecies variation

TITLE (TI): Direct Submission

L6 ANSWER 35 OF 58 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Extensive DNA inversions in the B. fragilis genome
control variable gene expression

TITLE (TI): Direct Submission

L6 ANSWER 36 OF 58 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): The genome of the heartwater agent Ehrlichia
ruminantium contains multiple tandem repeats of
actively variable copy number

TITLE (TI): Direct Submission

L6 ANSWER 37 OF 58 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): The complete genome sequence of Francisella tularensis,
the causative agent of tularemia

TITLE (TI): Direct Submission

L6 ANSWER 38 OF 58 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Complete genome sequence of Yersinia pestis strain
91001, an isolate avirulent to humans

TITLE (TI): Genetics of metabolic variations between Yersinia
pestis biovars and the proposal of a new biovar,
microtus

TITLE (TI): Direct Submission

L6 ANSWER 39 OF 58 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Extensive mosaic structure revealed by the complete
genome sequence of uropathogenic Escherichia coli

TITLE (TI): Direct Submission

L6 ANSWER 40 OF 58 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): The genome sequence of Bifidobacterium longum reflects its adaptation to the human gastrointestinal tract

TITLE (TI): Direct Submission

TITLE (TI): Direct Submission

L6 ANSWER 41 OF 58 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Genome sequence of Streptococcus mutans UA159, a cariogenic dental pathogen

TITLE (TI): Direct Submission

L6 ANSWER 42 OF 58 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Genome Sequence of Yersinia pestis KIM

TITLE (TI): Direct Submission

L6 ANSWER 43 OF 58 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Genome sequence of Yersinia pestis, the causative agent of plague

TITLE (TI): Annotation and evolutionary relationships of a small regulatory RNA gene micF and its target ompF in Yersinia species

TITLE (TI): Direct Submission

L6 ANSWER 44 OF 58 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): The genome sequence of the food-borne pathogen Campylobacter jejuni reveals hypervariable sequences

TITLE (TI): Re-annotation of Campylobacter jejuni NCTC11168

TITLE (TI): Direct Submission

TITLE (TI): Direct Submission

L6 ANSWER 45 OF 58 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Complete genome sequence of Clostridium perfringens, an anaerobic flesh-eater

TITLE (TI): Direct Submission

L6 ANSWER 46 OF 58 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Complete nucleotide sequence of the prophage VT2-Sakai carrying the verotoxin 2 genes of the enterohemorrhagic Escherichia coli O157:H7 derived from the Sakai outbreak

TITLE (TI): Comparative analysis of the whole set of rRNA operons between an enterohemorrhagic Escherichia coli O157:H7 Sakai strain and an Escherichia coli K-12 strain MG1655

TITLE (TI): Complete nucleotide sequence of the prophage VT1-Sakai carrying the Shiga toxin 1 genes of the enterohemorrhagic Escherichia coli O157:H7 strain derived from the Sakai outbreak

TITLE (TI): Complete genome sequence of enterohemorrhagic Escherichia coli O157:H7 and genomic comparison with a laboratory strain K-12

TITLE (TI): Direct Submission

L6 ANSWER 47 OF 58 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Genomic plasticity of the causative agent of melioidosis, Burkholderia pseudomallei

TITLE (TI): Direct Submission

L6 ANSWER 48 OF 58 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Insights into the evolution of *Yersinia pestis* through whole-genome comparison with *Yersinia pseudotuberculosis*

TITLE (TI): Direct Submission

L6 ANSWER 49 OF 58 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): The genome sequence of the enterobacterial phytopathogen *Erwinia carotovora* subsp. *atroseptica* SCRI1043 and functional genomic identification of novel virulence factors

TITLE (TI): Direct Submission

L6 ANSWER 50 OF 58 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Complete genomes of two clinical *Staphylococcus aureus* strains: evidence for the rapid evolution of virulence and drug resistance

TITLE (TI): Direct Submission

L6 ANSWER 51 OF 58 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Deciphering the biology of *Mycobacterium tuberculosis* from the complete genome sequence

TITLE (TI): Re-annotation of the genome sequence of *Mycobacterium tuberculosis* H37Rv

TITLE (TI): Direct Submission

L6 ANSWER 52 OF 58 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): The complete genome sequence and analysis of *Corynebacterium diphtheriae* NCTC13129

TITLE (TI): Direct Submission

L6 ANSWER 53 OF 58 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Comparative analysis of the genome sequences of *Bordetella pertussis*, *Bordetella parapertussis* and *Bordetella bronchiseptica*

TITLE (TI): Direct Submission

L6 ANSWER 54 OF 58 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): The complete genome sequence of *Mycobacterium bovis*

TITLE (TI): Direct Submission

L6 ANSWER 55 OF 58 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Genome Sequence and Comparative Analysis of the Solvent-Producing Bacterium *Clostridium acetobutylicum*

TITLE (TI): Direct Submission

L6 ANSWER 56 OF 58 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Complete genome sequence of the model actinomycete *Streptomyces coelicolor* A3(2)

TITLE (TI): Direct Submission

L6 ANSWER 57 OF 58 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Complete genome sequence of the model actinomycete *Streptomyces coelicolor* A3(2)

TITLE (TI): Direct Submission

L6 ANSWER 58 OF 58 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Genome sequence of enterohaemorrhagic Escherichia coli
O157:H7

TITLE (TI): Direct Submission

=> s l1(s) (TUBERCULOS? OR SMEGMAT?)
L7 172 L1(S) (TUBERCULOS? OR SMEGMAT?)

=> s l7 (s) (vector? or plasmid?)
L8 83 L7 (S) (VECTOR? OR PLASMID?)

=> s l8(s) (gpm?)
L9 17 L8(S) (GPM?)

=> dup rem l9
DUPLICATE IS NOT AVAILABLE IN 'GENBANK'.
ANSWERS FROM THESE FILES WILL BE CONSIDERED UNIQUE
PROCESSING COMPLETED FOR L9
L10 17 DUP REM L9 (0 DUPLICATES REMOVED)

=> d ti l10 1-17

L10 ANSWER 1 OF 17 USPATFULL on STN
TI Recombinant mycobacteria overexpressing D-alanine ligase gene and uses
therefore

L10 ANSWER 2 OF 17 USPATFULL on STN
TI Streptococcus pneumoniae polynucleotides and sequences

L10 ANSWER 3 OF 17 USPATFULL on STN
TI Nucleic acid and amino acid sequences relating to Streptococcus
pneumoniae for diagnostics and therapeutics

L10 ANSWER 4 OF 17 USPATFULL on STN
TI STREPTOCOCCUS PNEUMONIAE POLYNUCLEOTIDES AND SEQUENCES

L10 ANSWER 5 OF 17 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): The complete genome sequence of the European
Francisella tularensis subspecies tularensis isolate
FSC 198 suggests that it is derived from the archetypal
laboratory strain Schu S4, originally isolated in North
America

TITLE (TI): Direct Submission

L10 ANSWER 6 OF 17 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Complete sequence of chromosome of Mycobacterium sp.
MCS

TITLE (TI): Direct Submission

L10 ANSWER 7 OF 17 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): The partitioned Rhizobium etli genome: Genetic and
metabolic redundancy in seven interacting replicons

TITLE (TI): Direct Submission

L10 ANSWER 8 OF 17 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): The Chlamydophila abortus genome sequence reveals an
array of variable proteins that contribute to

interspecies variation
 TITLE (TI): The Chlamydomophila abortus genome sequence reveals an array of variable proteins that contribute to interspecies variation
 TITLE (TI): Direct Submission

L10 ANSWER 9 OF 17 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Extensive DNA inversions in the B. fragilis genome control variable gene expression
 TITLE (TI): Direct Submission

L10 ANSWER 10 OF 17 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Complete genome sequence of Yersinia pestis strain 91001, an isolate avirulent to humans
 TITLE (TI): Genetics of metabolic variations between Yersinia pestis biovars and the proposal of a new biovar, microtus
 TITLE (TI): Direct Submission

L10 ANSWER 11 OF 17 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Genome sequence of Streptococcus mutans UA159, a cariogenic dental pathogen
 TITLE (TI): Direct Submission

L10 ANSWER 12 OF 17 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Genome sequence of Yersinia pestis, the causative agent of plague
 TITLE (TI): Annotation and evolutionary relationships of a small regulatory RNA gene micF and its target ompF in Yersinia species
 TITLE (TI): Direct Submission

L10 ANSWER 13 OF 17 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Complete nucleotide sequence of the prophage VT2-Sakai carrying the verotoxin 2 genes of the enterohemorrhagic Escherichia coli O157:H7 derived from the Sakai outbreak
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 TITLE (TI): Complete genome sequence of enterohemorrhagic Escherichia coli O157:H7 and genomic comparison with a laboratory strain K-12
 TITLE (TI): Direct Submission

L10 ANSWER 14 OF 17 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Genomic plasticity of the causative agent of melioidosis, Burkholderia pseudomallei
 TITLE (TI): Direct Submission

L10 ANSWER 15 OF 17 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Insights into the evolution of Yersinia pestis through whole-genome comparison with Yersinia pseudotuberculosis

TITLE (TI): Direct Submission

L10 ANSWER 16 OF 17 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): The complete genome sequence and analysis of
Corynebacterium diphtheriae NCTC13129

TITLE (TI): Direct Submission

L10 ANSWER 17 OF 17 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Complete genome sequence of the model actinomycete
Streptomyces coelicolor A3(2)

TITLE (TI): Direct Submission

=> l8 and (gpm? or pbun?)

L8 IS NOT A RECOGNIZED COMMAND

The previous command name entered was not recognized by the system.

For a list of commands available to you in the current file, enter

"HELP COMMANDS" at an arrow prompt (=>).

=> s l8 and (gpm? or pbun?)

L11 38 L8 AND (GPM? OR PBUN?)

=> dup rem l11

DUPLICATE IS NOT AVAILABLE IN 'GENBANK'.

ANSWERS FROM THESE FILES WILL BE CONSIDERED UNIQUE

PROCESSING COMPLETED FOR L11

L12 38 DUP REM L11 (0 DUPLICATES REMOVED)

=> d ti l12 1-38

L12 ANSWER 1 OF 38 USPATFULL on STN

TI Novel polynucleotides

L12 ANSWER 2 OF 38 USPATFULL on STN

TI Nucleic acid and amino acid sequences relating to Enterobacter cloacae
for diagnostics and therapeutics

L12 ANSWER 3 OF 38 USPATFULL on STN

TI Nucleic acid and amino acid sequences relating to streptococcus
pneumoniae for diagnostics and therapeutics

L12 ANSWER 4 OF 38 USPATFULL on STN

TI Recombinant mycobacteria overexpressing D-alanine ligase gene and uses
therefore

L12 ANSWER 5 OF 38 USPATFULL on STN

TI Streptococcus pneumoniae polynucleotides and sequences

L12 ANSWER 6 OF 38 USPATFULL on STN

TI Novel human polynucleotides and polypeptides encoded thereby

L12 ANSWER 7 OF 38 USPATFULL on STN

TI Nucleic acid and amino acid sequences relating to Streptococcus
pneumoniae for diagnostics and therapeutics

L12 ANSWER 8 OF 38 USPATFULL on STN

TI Nucleic acid and amino acid sequences relating to Enterococcus faecalis
for diagnostics and therapeutics

L12 ANSWER 9 OF 38 USPATFULL on STN

TI Nucleic acid sequences and expression system relating to Enterococcus
faecium for diagnostics and therapeutics.

L12 ANSWER 10 OF 38 USPATFULL on STN
 TI Nucleic acid and amino acid sequences relating to *Acinetobacter baumannii* for diagnostics and therapeutics

L12 ANSWER 11 OF 38 USPATFULL on STN
 TI Novel Polynucleotides

L12 ANSWER 12 OF 38 USPATFULL on STN
 TI ENTEROCOCCUS FAECALIS POLYNUCLEOTIDES AND POLYPEPTIDES

L12 ANSWER 13 OF 38 USPATFULL on STN
 TI STREPTOCOCCUS PNEUMONIAE POLYNUCLEOTIDES AND SEQUENCES

L12 ANSWER 14 OF 38 GENBANK® COPYRIGHT 2006 on STN
 TITLE (TI): The genome of *Rhizobium leguminosarum* has recognizable core and accessory components
 TITLE (TI): Direct Submission

L12 ANSWER 15 OF 38 GENBANK® COPYRIGHT 2006 on STN
 TITLE (TI): The complete genome sequence of the European *Francisella tularensis* subspecies *tularensis* isolate FSC 198 suggests that it is derived from the archetypal laboratory strain Schu S4, originally isolated in North America
 TITLE (TI): Direct Submission

L12 ANSWER 16 OF 38 GENBANK® COPYRIGHT 2006 on STN
 TITLE (TI): Complete sequence of chromosome of *Mycobacterium* sp. MCS
 TITLE (TI): Direct Submission

L12 ANSWER 17 OF 38 GENBANK® COPYRIGHT 2006 on STN
 TITLE (TI): The partitioned *Rhizobium etli* genome: Genetic and metabolic redundancy in seven interacting replicons
 TITLE (TI): Direct Submission

L12 ANSWER 18 OF 38 GENBANK® COPYRIGHT 2006 on STN
 TITLE (TI): Complete Sequence of Chromosome 1 of *Rhodobacter sphaeroides* 2.4.1
 TITLE (TI): Direct Submission

L12 ANSWER 19 OF 38 GENBANK® COPYRIGHT 2006 on STN
 TITLE (TI): The *Chlamydophila abortus* genome sequence reveals an array of variable proteins that contribute to interspecies variation
 TITLE (TI): The *Chlamydophila abortus* genome sequence reveals an array of variable proteins that contribute to interspecies variation
 TITLE (TI): Direct Submission

L12 ANSWER 20 OF 38 GENBANK® COPYRIGHT 2006 on STN
 TITLE (TI): Extensive DNA inversions in the *B. fragilis* genome control variable gene expression
 TITLE (TI): Direct Submission

L12 ANSWER 21 OF 38 GENBANK® COPYRIGHT 2006 on STN
 TITLE (TI): The complete genome sequence of *Francisella tularensis*,

TITLE (TI): the causative agent of tularemia
 Direct Submission

L12 ANSWER 22 OF 38 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Complete genome sequence of Yersinia pestis strain 91001, an isolate avirulent to humans
 TITLE (TI): Genetics of metabolic variations between Yersinia pestis biovars and the proposal of a new biovar, microtus
 TITLE (TI): Direct Submission

L12 ANSWER 23 OF 38 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Genome sequence of Streptococcus mutans UA159, a cariogenic dental pathogen
 TITLE (TI): Direct Submission

L12 ANSWER 24 OF 38 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Genome Sequence of Yersinia pestis KIM
 TITLE (TI): Direct Submission

L12 ANSWER 25 OF 38 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Genome sequence of Yersinia pestis, the causative agent of plague
 TITLE (TI): Annotation and evolutionary relationships of a small regulatory RNA gene micF and its target ompF in Yersinia species
 TITLE (TI): Direct Submission

L12 ANSWER 26 OF 38 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): The genome sequence of the food-borne pathogen Campylobacter jejuni reveals hypervariable sequences
 TITLE (TI): Re-annotation of Campylobacter jejuni NCTC11168
 TITLE (TI): Direct Submission
 TITLE (TI): Direct Submission

L12 ANSWER 27 OF 38 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Complete genome sequence of Clostridium perfringens, an anaerobic flesh-eater
 TITLE (TI): Direct Submission

L12 ANSWER 28 OF 38 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Complete nucleotide sequence of the prophage VT2-Sakai carrying the verotoxin 2 genes of the enterohemorrhagic Escherichia coli O157:H7 derived from the Sakai outbreak
 TITLE (TI): Comparative analysis of the whole set of rRNA operons between an enterohemorrhagic Escherichia coli O157:H7 Sakai strain and an Escherichia coli K-12 strain MG1655
 TITLE (TI): Complete nucleotide sequence of the prophage VT1-Sakai carrying the Shiga toxin 1 genes of the enterohemorrhagic Escherichia coli O157:H7 strain derived from the Sakai outbreak
 TITLE (TI): Complete genome sequence of enterohemorrhagic Escherichia coli O157:H7 and genomic comparison with a laboratory strain K-12
 TITLE (TI): Direct Submission

L12 ANSWER 29 OF 38 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Genomic plasticity of the causative agent of
melioidosis, *Burkholderia pseudomallei*
TITLE (TI): Direct Submission

L12 ANSWER 30 OF 38 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Insights into the evolution of *Yersinia pestis* through
whole-genome comparison with *Yersinia*
pseudotuberculosis
TITLE (TI): Direct Submission

L12 ANSWER 31 OF 38 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): The genome sequence of the enterobacterial
phytopathogen *Erwinia carotovora* subsp. *atroseptica*
SCRI1043 and functional genomic identification of novel
virulence factors
TITLE (TI): Direct Submission

L12 ANSWER 32 OF 38 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Complete genomes of two clinical *Staphylococcus aureus*
strains: evidence for the rapid evolution of virulence
and drug resistance
TITLE (TI): Direct Submission

L12 ANSWER 33 OF 38 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Deciphering the biology of *Mycobacterium tuberculosis*
from the complete genome sequence
TITLE (TI): Re-annotation of the genome sequence of *Mycobacterium*
tuberculosis H37Rv
TITLE (TI): Direct Submission

L12 ANSWER 34 OF 38 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): The complete genome sequence and analysis of
Corynebacterium diphtheriae NCTC13129
TITLE (TI): Direct Submission

L12 ANSWER 35 OF 38 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Genome Sequence and Comparative Analysis of the
Solvent-Producing Bacterium *Clostridium acetobutylicum*
TITLE (TI): Direct Submission

L12 ANSWER 36 OF 38 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Complete genome sequence of the model actinomycete
Streptomyces coelicolor A3(2)
TITLE (TI): Direct Submission

L12 ANSWER 37 OF 38 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Complete genome sequence of the model actinomycete
Streptomyces coelicolor A3(2)
TITLE (TI): Direct Submission

L12 ANSWER 38 OF 38 GENBANK® COPYRIGHT 2006 on STN

TITLE (TI): Genome sequence of enterohaemorrhagic *Escherichia coli*
O157:H7
TITLE (TI): Direct Submission

=> d ibib abs 112 1-13

L12 ANSWER 1 OF 38 USPATFULL on STN

ACCESSION NUMBER: 2006:268028 USPATFULL
TITLE: Novel polynucleotides
INVENTOR(S): Nakagawa, Satoshi, Tokyo, JAPAN
Mizoguchi, Hiroshi, Tokyo, JAPAN
Ando, Seiko, Tokyo, JAPAN
Hayashi, Mikiro, Tokyo, JAPAN
Ochiai, Keiko, Tokyo, JAPAN
Yokoi, Haruhiko, Tokyo, JAPAN
Tateishi, Naoko, Tokyo, JAPAN
Senoh, Akihiro, Tokyo, JAPAN
Ikeda, Masato, Tokyo, JAPAN
Ozaki, Akio, Hofu-shi, JAPAN
PATENT ASSIGNEE(S): Kyowa Hakko Kogyo Co., Ltd., Tokyo, JAPAN (non-U.S. corporation)

	NUMBER	KIND	DATE
PATENT INFORMATION:	US 2006228712	A1	20061012
APPLICATION INFO.:	US 2004-805394	A1	20040322 (10)
RELATED APPLN. INFO.:	Division of Ser. No. US 2000-738626, filed on 18 Dec 2000, ABANDONED		

	NUMBER	DATE
PRIORITY INFORMATION:	JP 1999-377484	19991216
	JP 2000-159162	20000407
	JP 2000-280988	20000803
DOCUMENT TYPE:	Utility	
FILE SEGMENT:	APPLICATION	
LEGAL REPRESENTATIVE:	NIXON & VANDERHYE, PC, 901 NORTH GLEBE ROAD, 11TH FLOOR, ARLINGTON, VA, 22203, US	
NUMBER OF CLAIMS:	67	
EXEMPLARY CLAIM:	1	
NUMBER OF DRAWINGS:	4 Drawing Page(s)	
LINE COUNT:	14357	

AB Novel polynucleotides derived from microorganisms belonging to coryneform bacteria and fragments thereof, polypeptides encoded by the polynucleotides and fragments thereof, polynucleotide arrays comprising the polynucleotides and fragments thereof, recording media in which the nucleotide sequences of the polynucleotide and fragments thereof have been recorded which are readable in a computer, and use of them.

L12 ANSWER 2 OF 38 USPATFULL on STN

ACCESSION NUMBER: 2006:113827 USPATFULL
TITLE: Nucleic acid and amino acid sequences relating to Enterobacter cloacae for diagnostics and therapeutics
INVENTOR(S): Weinstock, Keith G., Westborough, MA, UNITED STATES
Deloughery, Craig, Medford, MA, UNITED STATES
Bush, David, Somerville, MA, UNITED STATES
PATENT ASSIGNEE(S): Genome Therapeutics Corporation, Waltham, MA, UNITED STATES (U.S. corporation)

	NUMBER	KIND	DATE
PATENT INFORMATION:	US 7041814	B1	20060509
APPLICATION INFO.:	US 1999-252691		19990218 (9)

	NUMBER	DATE
PRIORITY INFORMATION:	US 1998-94145P	19980724 (60)

US 1998-74787P 19980218 (60)
DOCUMENT TYPE: Utility
FILE SEGMENT: GRANTED
PRIMARY EXAMINER: Smith, Lynette R. F.
ASSISTANT EXAMINER: Portner, Ginny Allen
LEGAL REPRESENTATIVE: Buchanan Ingersoll PC
NUMBER OF CLAIMS: 9
EXEMPLARY CLAIM: 1
LINE COUNT: 19563

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The invention provides isolated polypeptide and nucleic acid sequences derived from *Enterobacter cloacae* that are useful in diagnosis and therapy of pathological conditions; antibodies against the polypeptides; and methods for the production of the polypeptides. The invention also provides methods for the detection, prevention and treatment of pathological conditions resulting from bacterial infection.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L12 ANSWER 3 OF 38 USPATFULL on STN

ACCESSION NUMBER: 2005:158196 USPATFULL
TITLE: Nucleic acid and amino acid sequences relating to *streptococcus pneumoniae* for diagnostics and therapeutics
INVENTOR(S): Doucette-Stamm, Lynn A., Framingham, MA, UNITED STATES
Bush, David, Somerville, MA, UNITED STATES

	NUMBER	KIND	DATE
PATENT INFORMATION:	US 2005136404	A1	20050623
APPLICATION INFO.:	US 2003-617320	A1	20030710 (10)
RELATED APPLN. INFO.:	Division of Ser. No. US 1998-107433, filed on 30 Jun 1998, PENDING		

	NUMBER	DATE
PRIORITY INFORMATION:	US 1997-51553P	19970702 (60)
	US 1998-85131P	19980512 (60)
DOCUMENT TYPE:	Utility	
FILE SEGMENT:	APPLICATION	
LEGAL REPRESENTATIVE:	Robert L. Spadafora, Genome Therapeutics Corporation, 100 Beaver Street, Waltham, MA, 02453, US	
NUMBER OF CLAIMS:	28	
EXEMPLARY CLAIM:	1	
LINE COUNT:	12957	

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The invention provides isolated polypeptide and nucleic acid sequences derived from *Streptococcus pneumonia* that are useful in diagnosis and therapy of pathological conditions; antibodies against the polypeptides; and methods for the production of the polypeptides. The invention also provides methods for the detection, prevention and treatment of pathological conditions resulting from bacterial infection.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L12 ANSWER 4 OF 38 USPATFULL on STN

ACCESSION NUMBER: 2004:307159 USPATFULL
TITLE: Recombinant mycobacteria overexpressing D-alanine ligase gene and uses therefore
INVENTOR(S): Barletta, Raul G., Lincoln, NE, UNITED STATES
Feng, Zhengyu, Austin, TX, UNITED STATES
PATENT ASSIGNEE(S): The Board of Regents, University of Nebraska-Lincoln (U.S. corporation)

NUMBER	KIND	DATE
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PATENT INFORMATION:	US 2004241830	A1	20041202	
APPLICATION INFO.:	US 2003-738938	A1	20031217	(10)

NUMBER	DATE
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PRIORITY INFORMATION:	US 2002-434200P	20021217 (60)
DOCUMENT TYPE:	Utility	
FILE SEGMENT:	APPLICATION	
LEGAL REPRESENTATIVE:	STINSON MORRISON HECKER LLP, ATTN: PATENT GROUP, 1201 WALNUT STREET, SUITE 2800, KANSAS CITY, MO, 64106-2150	
NUMBER OF CLAIMS:	23	
EXEMPLARY CLAIM:	1	
NUMBER OF DRAWINGS:	7 Drawing Page(s)	
LINE COUNT:	1431	

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB Recombinant mycobacterial strains which overproduce essential biosynthetic enzymes of pathogenic mycobacteria are provided. These strains overproduce enzymes involved in the synthesis and incorporation of D-alanine into mycobacterial peptidoglycan, the backbone of the mycobacterial cell wall. These overproducing strains may be used as reference strains in in vitro screening methods to identify antimycobacterial agents.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L12 ANSWER 5 OF 38 USPATFULL on STN

ACCESSION NUMBER:	2004:38579 USPATFULL
TITLE:	Streptococcus pneumoniae polynucleotides and sequences
INVENTOR(S):	Kunsch, Charles A., Norcross, GA, UNITED STATES Choi, Gil H., Rockville, MD, UNITED STATES Dillon, Patrick J., Carlsbad, CA, UNITED STATES Rosen, Craig A., Laytonsville, MD, UNITED STATES Barash, Steven C., Rockville, MD, UNITED STATES Fannon, Michael R., Silver Spring, MD, UNITED STATES Dougherty, Brian A., Killingworth, CT, UNITED STATES
PATENT ASSIGNEE(S):	Human Genome Sciences, Inc., Rockville, MD, UNITED STATES, 20850 (U.S. corporation)

NUMBER	KIND	DATE
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PATENT INFORMATION:	US 2004029118	A1	20040212
APPLICATION INFO.:	US 2002-158844	A1	20020603 (10)
RELATED APPLN. INFO.:	Division of Ser. No. US 1997-961527, filed on 30 Oct 1997, GRANTED, Pat. No. US 6420135		

NUMBER	DATE
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PRIORITY INFORMATION:	US 1996-29960P	19961031 (60)
DOCUMENT TYPE:	Utility	
FILE SEGMENT:	APPLICATION	
LEGAL REPRESENTATIVE:	HUMAN GENOME SCIENCES INC, 9410 KEY WEST AVENUE, ROCKVILLE, MD, 20850	
NUMBER OF CLAIMS:	20	
EXEMPLARY CLAIM:	1	
NUMBER OF DRAWINGS:	2 Drawing Page(s)	
LINE COUNT:	9165	

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The present invention provides polynucleotide sequences of the genome of Streptococcus pneumoniae, polypeptide sequences encoded by the polynucleotide sequences, corresponding polynucleotides and polypeptides, vectors and hosts comprising the polynucleotides, and assays and other uses thereof. The present invention further provides polynucleotide and polypeptide sequence information stored on computer readable media, and computer-based systems and methods which facilitate

its use.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L12 ANSWER 6 OF 38 USPATFULL on STN

ACCESSION NUMBER: 2004:12955 USPATFULL
TITLE: Novel human polynucleotides and polypeptides encoded thereby
INVENTOR(S): Leach, Martin D., Madison, CT, UNITED STATES
Shimkets, Richard A., Guilford, CT, UNITED STATES

	NUMBER	KIND	DATE
PATENT INFORMATION:	US 2004009474	A1	20040115
APPLICATION INFO.:	US 2001-864408	A1	20010524 (9)

	NUMBER	DATE
PRIORITY INFORMATION:	US 2000-206690P	20000524 (60)
DOCUMENT TYPE:	Utility	
FILE SEGMENT:	APPLICATION	
LEGAL REPRESENTATIVE:	Ivor R. Elrifi, Esq., MIntz, Levin, Cohn, Ferris,, Glovsky and Popeo, P.C., One Financial Center, Boston, MA, 02111	
NUMBER OF CLAIMS:	32	
EXEMPLARY CLAIM:	1	
LINE COUNT:	21366	

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The present invention provides ORFX, a novel isolated polypeptide, as well as a polynucleotide encoding ORFX and antibodies that immunospecifically bind to ORFX or any derivative, variant, mutant, or fragment of the ORFX polypeptide, polynucleotide or antibody. The invention additionally provides methods in which the ORFX polypeptide, polynucleotide and antibody are used in detection and treatment of a broad range of pathological states, as well as to others uses.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L12 ANSWER 7 OF 38 USPATFULL on STN

ACCESSION NUMBER: 2004:250212 USPATFULL
TITLE: Nucleic acid and amino acid sequences relating to Streptococcus pneumoniae for diagnostics and therapeutics
INVENTOR(S): Doucette-Stamm, Lynn A., Framingham, MA, United States
Bush, David, Somerville, MA, United States
PATENT ASSIGNEE(S): Genome Therapeutics Corporation, Waltham, MA, United States (U.S. corporation)

	NUMBER	KIND	DATE
PATENT INFORMATION:	US 6800744	B1	20041005
APPLICATION INFO.:	US 1998-107433		19980630 (9)

	NUMBER	DATE
PRIORITY INFORMATION:	US 1998-85131P	19980512 (60)
	US 1997-51553P	19970702 (60)
DOCUMENT TYPE:	Utility	
FILE SEGMENT:	GRANTED	
PRIMARY EXAMINER:	Brusca, John S.	
ASSISTANT EXAMINER:	Zhou, Shubo "Joe "	
LEGAL REPRESENTATIVE:	Genome Therapeutics Corporation	
NUMBER OF CLAIMS:	14	
EXEMPLARY CLAIM:	1	
NUMBER OF DRAWINGS:	0 Drawing Figure(s); 0 Drawing Page(s)	

LINE COUNT: 11545

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The invention provides isolated polypeptide and nucleic acid sequences derived from Streptococcus pneumonia that are useful in diagnosis and therapy of pathological conditions; antibodies against the polypeptides; and methods for the production of the polypeptides. The invention also provides methods for the detection, prevention and treatment of pathological conditions resulting from bacterial infection.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L12 ANSWER 8 OF 38 USPATFULL on STN

ACCESSION NUMBER: 2003:240330 USPATFULL

TITLE: Nucleic acid and amino acid sequences relating to Enterococcus faecalis for diagnostics and therapeutics

INVENTOR(S): Doucette-Stamm, Lynn A., 14 Flanagan Dr., Framingham, MA, United States 01701
Bush, David, 205 Holland St., Somerville, MA, United States 02144

	NUMBER	KIND	DATE
PATENT INFORMATION:	US 6617156	B1	20030909
APPLICATION INFO.:	US 1998-134000		19980813 (9)

	NUMBER	DATE
PRIORITY INFORMATION:	US 1997-55778P	19970815 (60)
DOCUMENT TYPE:	Utility	
FILE SEGMENT:	GRANTED	
PRIMARY EXAMINER:	Mosher, Mary E.	
LEGAL REPRESENTATIVE:	Genome Therapeutics Corporation	
NUMBER OF CLAIMS:	19	
EXEMPLARY CLAIM:	1,5,14	
NUMBER OF DRAWINGS:	0 Drawing Figure(s); 0 Drawing Page(s)	
LINE COUNT:	13738	

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The invention provides isolated polypeptide and nucleic acid sequences derived from Enterococcus faecalis that are useful in diagnosis and therapy of pathological conditions; antibodies against the polypeptides; and methods for the production of the polypeptides. The invention also provides methods for the detection, prevention and treatment of pathological conditions resulting from bacterial infection.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L12 ANSWER 9 OF 38 USPATFULL on STN

ACCESSION NUMBER: 2003:169096 USPATFULL

TITLE: Nucleic acid sequences and expression system relating to Enterococcus faecium for diagnostics and therapeutics

INVENTOR(S): Doucette-Stamm, Lynn A., Framingham, MA, United States
Bush, David, Somerville, MA, United States

PATENT ASSIGNEE(S): Genome Therapeutics Corporation, Waltham, MA, United States (U.S. corporation)

	NUMBER	KIND	DATE
PATENT INFORMATION:	US 6583275	B1	20030624
APPLICATION INFO.:	US 1998-107532		19980630 (9)

	NUMBER	DATE
PRIORITY INFORMATION:	US 1998-85598P	19980514 (60)
	US 1997-51571P	19970702 (60)

DOCUMENT TYPE: Utility
FILE SEGMENT: GRANTED
PRIMARY EXAMINER: Marschel, Ardin H.
LEGAL REPRESENTATIVE: Genome Therapeutics Corporation
NUMBER OF CLAIMS: 34
EXEMPLARY CLAIM: 1
NUMBER OF DRAWINGS: 0 Drawing Figure(s); 0 Drawing Page(s)
LINE COUNT: 15265

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The invention provides isolated polypeptide and nucleic acid sequences derived *Enterococcus faecium* that are useful in diagnosis and therapy of pathological conditions; antibodies against the polypeptides; and methods for the production of the polypeptides. The invention also provides methods for the detection, prevention and treatment of pathological conditions resulting from bacterial infection.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L12 ANSWER 10 OF 38 USPATFULL on STN

ACCESSION NUMBER: 2003:130010 USPATFULL
TITLE: Nucleic acid and amino acid sequences relating to *Acinetobacter baumannii* for diagnostics and therapeutics
INVENTOR(S): Breton, Gary, Marlborough, MA, United States
Bush, David, Somerville, MA, United States
PATENT ASSIGNEE(S): Genome Therapeutics Corporation, Waltham, MA, United States (U.S. corporation)

	NUMBER	KIND	DATE
PATENT INFORMATION:	US 6562958	B1	20030513
APPLICATION INFO.:	US 1999-328352		19990604 (9)

	NUMBER	DATE
PRIORITY INFORMATION:	US 1998-88701P	19980609 (60)
DOCUMENT TYPE:	Utility	
FILE SEGMENT:	GRANTED	
PRIMARY EXAMINER:	Borin, Michael	
LEGAL REPRESENTATIVE:	Genome Therapeutics Corporation	
NUMBER OF CLAIMS:	15	
EXEMPLARY CLAIM:	1	
NUMBER OF DRAWINGS:	0 Drawing Figure(s); 0 Drawing Page(s)	
LINE COUNT:	16618	

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The invention provides isolated polypeptide and nucleic acid sequences derived from *Acinetobacter mirabilis* that are useful in diagnosis and therapy of pathological conditions; antibodies against the polypeptides; and methods for the production of the polypeptides. The invention also provides methods for the detection, prevention and treatment of pathological conditions resulting from bacterial infection.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L12 ANSWER 11 OF 38 USPATFULL on STN

ACCESSION NUMBER: 2002:343879 USPATFULL
TITLE: Novel Polynucleotides
INVENTOR(S): Nakagawa, Satoshi, Tokyo, JAPAN
Mizoguchi, Hiroshi, Tokyo, JAPAN
Ando, Seiko, Tokyo, JAPAN
Hayashi, Mikiro, Tokyo, JAPAN
Ochiai, Keiko, Tokyo, JAPAN
Yokoi, Haruhiko, Tokyo, JAPAN
Tateishi, Naoko, Tokyo, JAPAN
Senoh, Akihiro, Tokyo, JAPAN

Ikeda, Masato, Tokyo, JAPAN
Ozaki, Akio, Hofu-shi, JAPAN

	NUMBER	KIND	DATE
PATENT INFORMATION:	US 2002197605	A1	20021226
APPLICATION INFO.:	US 2000-738626	A1	20001218 (9)

	NUMBER	DATE
PRIORITY INFORMATION:	JP 1999-377484	19991216
	JP 2000-159162	20000407
	JP 2000-280988	20000803
DOCUMENT TYPE:	Utility	
FILE SEGMENT:	APPLICATION	
LEGAL REPRESENTATIVE:	NIXON & VANDERHYE P.C., 8th Floor, 1100 North Glebe Road, Arlington, VA, 22201	
NUMBER OF CLAIMS:	68	
EXEMPLARY CLAIM:	1	
NUMBER OF DRAWINGS:	4 Drawing Page(s)	
LINE COUNT:	13673	

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB Novel polynucleotides derived from microorganisms belonging to coryneform bacteria and fragments thereof, polypeptides encoded by the polynucleotides and fragments thereof, polynucleotide arrays comprising the polynucleotides and fragments thereof, recording media in which the nucleotide sequences of the polynucleotide and fragments thereof have been recorded which are readable in a computer, and use of them.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L12 ANSWER 12 OF 38 USPATFULL on STN

ACCESSION NUMBER: 2002:221971 USPATFULL
TITLE: ENTEROCOCCUS FAECALIS POLYNUCLEOTIDES AND POLYPEPTIDES
INVENTOR(S): KUNSCH, CHARLES A., ATLANTA, GA, UNITED STATES
DILLON, PATRICK J., CARLSBAD, CA, UNITED STATES
BARASH, STEVEN, ROCKVILLE, MD, UNITED STATES

	NUMBER	KIND	DATE
PATENT INFORMATION:	US 2002120116	A1	20020829
APPLICATION INFO.:	US 1998-70927	A1	19980504 (9)
DOCUMENT TYPE:	Utility		
FILE SEGMENT:	APPLICATION		
LEGAL REPRESENTATIVE:	HUMAN GENOME SCIENCES INC, 9410 KEY WEST AVENUE, ROCKVILLE, MD, 20850		
NUMBER OF CLAIMS:	18		
EXEMPLARY CLAIM:	1		
NUMBER OF DRAWINGS:	2 Drawing Page(s)		
LINE COUNT:	13315		

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The present invention provides polynucleotide sequences of the genome of Enterococcus faecalis, polypeptide sequences encoded by the polynucleotide sequences, corresponding polynucleotides and polypeptides, vectors and hosts comprising the polynucleotides, and assays and other uses thereof. The present invention further provides polynucleotide and polypeptide sequence information stored on computer readable media, and computer-based systems and methods which facilitate its use.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L12 ANSWER 13 OF 38 USPATFULL on STN

ACCESSION NUMBER: 2002:55159 USPATFULL
TITLE: STREPTOCOCCUS PNEUMONIAE POLYNUCLEOTIDES AND SEQUENCES

INVENTOR(S) : KUNSCH, CHARLES A., GAITHERSBURG, MD, UNITED STATES
 CHOI, GIL H., ROCKVILLE, MD, UNITED STATES
 DILLON, PATRICK J., CARLSBAD, CA, UNITED STATES
 ROSEN, CRAIG A., LAYTONSVILLE, MD, UNITED STATES
 BARASH, STEVEN C., ROCKVILLE, MD, UNITED STATES
 FANNON, MICHAEL R., SILVER SPRING, MD, UNITED STATES
 DOUGHERTY, BRIAN A., MT. AIRY, MD, UNITED STATES

	NUMBER	KIND	DATE
PATENT INFORMATION:	US 2002032323	A1	20020314
	US 6420135	B2	20020716
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	NUMBER	DATE
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LEGAL REPRESENTATIVE:	HUMAN GENOME SCIENCES INC, 9410 KEY WEST AVENUE, ROCKVILLE, MD, 20850	
NUMBER OF CLAIMS:	20	
EXEMPLARY CLAIM:	1	
NUMBER OF DRAWINGS:	2 Drawing Page(s)	
LINE COUNT:	7752	

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The present invention provides polynucleotide sequences of the genome of Streptococcus pneumoniae, polypeptide sequences encoded by the polynucleotide sequences, corresponding polynucleotides and polypeptides, vectors and hosts comprising the polynucleotides, and assays and other uses thereof. The present invention further provides polynucleotide and polypeptide sequence information stored on computer readable media, and computer-based systems and methods which facilitate its use.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

=> d his full

(FILE 'HOME' ENTERED AT 13:53:33 ON 12 OCT 2006)

INDEX 'ADISCTI, ADISINSIGHT, ADISNEWS, AGRICOLA, ANABSTR, ANTE, AQUALINE, AQUASCI, BIOENG, BIOSIS, BIOTECHABS, BIOTECHDS, BIOTECHNO, CABA, CAPLUS, CEABA-VTB, CIN, CONFSCI, CROPB, CROPB, DDFB, DDFU, DGENE, DISSABS, DRUGB, DRUGMONOG2, DRUGU, EMBAL, EMBASE, ...' ENTERED AT 13:54:08 ON 12 OCT 2006
 SEA DDL? OR (ALANI?(S)LIGAS?)

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18  FILE ADISCTI
1   FILE ADISINSIGHT
30  FILE AGRICOLA
13  FILE ANABSTR
7   FILE ANTE
2   FILE AQUALINE
3   FILE AQUASCI
51  FILE BIOENG
654 FILE BIOSIS
72  FILE BIOTECHABS
72  FILE BIOTECHDS
250 FILE BIOTECHNO
105 FILE CABA
715 FILE CAPLUS
31  FILE CEABA-VTB
16  FILE CIN
17  FILE CONFSCI

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17 FILE CROPU
 6 FILE DDFB
 72 FILE DDFU
 594 FILE DGENE
 79 FILE DISSABS
 6 FILE DRUGB
 1 FILE DRUGMONOG2
 104 FILE DRUGU
 15 FILE EMBAL
 594 FILE EMBASE
 344 FILE ESBIODBASE
 6 FILE FROSTI
 13 FILE FSTA
 1795 FILE GENBANK
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 258 FILE IFIPAT
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 81 FILE JICST-EPLUS
 369 FILE LIFESCI
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 701 FILE PROMT
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 10 FILE RDISCLOSURE
 808 FILE SCISEARCH
 321 FILE TOXCENTER
 2690 FILE USPATFULL
 308 FILE USPAT2
 1 FILE VETU
 5 FILE WATER
 185 FILE WPIDS
 1 FILE WPIFV
 185 FILE WPINDEX
 11 FILE IPA
 234 FILE NLDB

L1 QUE DDL? OR (ALANI?(S) LIGAS?)

 D RANK

FILE 'USPATFULL, GENBANK, SCISEARCH, CAPLUS, PROMT, BIOSIS, PASCAL, EMBASE, MEDLINE, LIFESCI' ENTERED AT 13:56:02 ON 12 OCT 2006

L2 9522 SEA DDL? OR (ALANI?(S) LIGAS?)
 L3 501 SEA L2 AND (TUBERCULOS? OR SMEGMAT?)
 L4 399 SEA L3 AND (VECTOR? OR PLASMID?)
 L5 58 SEA L4 AND GPM?
 L6 58 DUP REM L5 (0 DUPLICATES REMOVED)
 D TI L6 1-58
 L7 172 SEA L1(S) (TUBERCULOS? OR SMEGMAT?)
 L8 83 SEA L7 (S) (VECTOR? OR PLASMID?)
 L9 17 SEA L8(S) (GPM?)
 L10 17 DUP REM L9 (0 DUPLICATES REMOVED)
 D TI L10 1-17
 L11 38 SEA L8 AND (GPM? OR PBUN?)
 L12 38 DUP REM L11 (0 DUPLICATES REMOVED)
 D TI L12 1-38
 D IBIB ABS L12 1-13

FILE HOME



Search



for



NiceZyme View of ENZYME: EC 6.3.2.4

Official Name

D-alanine--D-alanine ligase.

Alternative Name(s)

Alanine:alanine ligase (ADP-forming).

Alanylalanine synthetase.

D-Ala-D-Ala synthetase.

D-alanyl-D-alanine synthetase.

D-alanylalanine synthetase.

Reaction catalysed

ATP + 2 D-alanine <=> ADP + phosphate + D-alanyl-D-alanine

Comment(s)

Involved with EC 6.3.2.7 or EC 6.3.2.13, EC 6.3.2.8, EC 6.3.2.9 and EC 6.3.2.10 in the synthesis of a cell-wall peptide.

Cross-references

Biochemical Pathways; map number(s)

N3

PROSITE

PDOC00659

BRENDA

6.3.2.4

PUMA2

6.3.2.4

PRIAM enzyme-specific profiles

6.3.2.4

KEGG Ligand Database for Enzyme Nomenclature

6.3.2.4

IUBMB Enzyme Nomenclature

6.3.2.4

IntEnz

6.3.2.4

MEDLINE

Find literature relating to 6.3.2.4

MetaCyc

6.3.2.4

Q81D33, DDLA_BACCR;	Q89GA7, DDLA_BRAJA;	Q8YHR9, DDLA_BRUME;
Q8G044, DDLA_BRUSU;	Q7NV72, DDLA_CHRVO;	Q8XKS9, DDLA_CLOPE;
Q897P8, DDLA_CLOTE;	P0A6J9, DDLA_ECO57;	Q8FKE3, DDLA_ECOL6;
P0A6J8, DDLA_ECOLI;	Q9HWI0, DDLA_PSEAE;	Q88EV6, DDLA_PSEPK;
Q87XJ6, DDLA_PSESM;	Q98JS0, DDLA_RHILO;	P0A1F1, DDLA_SALTI;
P0A1F0, DDLA_SALTY;	P0A6K0, DDLA_SHIFL;	Q8PDW3, DDLA_XANCP;
Q81IU1, DDLB_BACCR;	Q89PS5, DDLB_BRAJA;	Q8YI63, DDLB_BRUME;
Q8FZP5, DDLB_BRUSU;	Q7NQ01, DDLB_CHRVO;	Q8XM71, DDLB_CLOPE;
Q898Z5, DDLB_CLOTE;	Q8X9Y6, DDLB_ECO57;	Q8FL63, DDLB_ECOL6;
P07862, DDLB_ECOLI;	Q9LCT6, DDLB_PSEAE;	Q88N74, DDLB_PSEPK;

Q87WY7, DDLB_PSESM;	Q98KB6, DDLB_RHILO;	Q8Z9G7, DDLB_SALTI;
Q8ZRU1, DDLB_SALTY;	Q83MF7, DDLB_SHIFL;	Q8PCJ8, DDLB_XANCP;
Q8UDN3, DDL_AGRT5;	P35660, DDL_ANACE;	Q8YY71, DDL_ANASP;
O66806, DDL_AQUAE;	Q81Q29, DDL_BACAN;	Q9KCF0, DDL_BACHD;
P96612, DDL_BACSU;	Q8A1F3, DDL_BACTN;	Q8G7C4, DDL_BIFLO;
Q7VRX1, DDL_BLOFL;	Q7WFS4, DDL_BORBR;	O51218, DDL_BORBU;
Q7W4B6, DDL_BORPA;	Q7VUQ5, DDL_BORPE;	O51927, DDL_BUCAP;
P59435, DDL_BUCBP;	Q9PPC2, DDL_CAMJE;	Q9A5A9, DDL_CAUCR;
Q8KCR8, DDL_CHLTE;	Q97F58, DDL_CLOAB;	Q8FPQ9, DDL_COREF;
Q8NQV2, DDL_CORGL;	Q83BZ9, DDL_COXBU;	Q9RXF1, DDL_DEIRA;
Q47758, DDL_ENTFA;	Q47823, DDL_ENTGA;	Q47827, DDL_ENTHR;
Q8RDQ4, DDL_FUSNN;	Q7MNN1, DDL_GLOVI;	Q7VMY2, DDL_HAEDU;
Q4QLF6, DDL_HAEI8;	P44405, DDL_HAEIN;	Q7VJW2, DDL_HELHP;
Q9ZLA5, DDL_HELPJ;	P56191, DDL_HELPY;	Q9CIL5, DDL_LACLA;
Q88UV8, DDL_LACPL;	Q72R93, DDL_LEPIC;	Q8F4I2, DDL_LEPIN;
Q48745, DDL_LEUME;	Q92DG5, DDL_LISIN;	Q721W2, DDL_LISMF;
Q8Y8P1, DDL_LISMO;	Q65RY9, DDL_MANSM;	Q7TXH9, DDL_MYCBO;
Q9CBS0, DDL_MYCLE;	Q9ZGN0, DDL_MYCSM;	P95114, DDL_MYCTU;
Q9JSZ9, DDL_NEIMA;	Q9K0Y0, DDL_NEIMB;	Q82VS0, DDL_NITEU;
Q8ERJ6, DDL_OCEIH;	P57819, DDL_PASMU;	Q7N149, DDL_PHOLL;
Q7MWA2, DDL_PORGI;	Q7VAS4, DDL_PROMA;	Q7V8L9, DDL_PROMM;
Q7V0F6, DDL_PROMP;	Q8XVI9, DDL_RALSO;	Q92NM4, DDL_RHIME;
Q92IT7, DDL_RICCN;	Q9ZDS6, DDL_RICPR;	Q8EEZ2, DDL_SHEON;
Q5HEB7, DDL_STAAC;	P63891, DDL_STAAM;	P63892, DDL_STAAN;
Q6GEZ1, DDL_STAAR;	Q6G7M7, DDL_STAAS;	Q8NVH8, DDL_STAAB;
Q5HMD8, DDL_STAEQ;	Q8CRP5, DDL_STAES;	Q8E640, DDL_STRA3;
Q8E0G6, DDL_STRA5;	Q82JS5, DDL_STRAW;	Q9ZBR9, DDL_STRCO;
P95803, DDL_STRMU;	Q99Z34, DDL_STRP1;	Q8K6X7, DDL_STRP3;
Q5XB93, DDL_STRP6;	Q8P0C4, DDL_STRP8;	O54631, DDL_STRPN;
Q8DNV5, DDL_STRR6;	Q8DLH0, DDL_SYNEL;	Q7U5Q9, DDL_SYNPX;
P73632, DDL_SYNY3;	P46805, DDL_THEMA;	Q8R778, DDL_THETN;
O83676, DDL_TREPA;	Q83I36, DDL_TROW8;	Q83G28, DDL_TROWT;
Q9KM17, DDL_VIBCH;	Q87HS0, DDL_VIBPA;	Q8D781, DDL_VIBVU;
Q7ME85, DDL_VIBVY;	Q8D2Y5, DDL_WIGBR;	Q7MA71, DDL_WOLSU;
Q8PPA6, DDL_XANAC;	Q9PF79, DDL_XYLFA;	Q87AG1, DDL_XYLFT;
Q8ZIE7, DDL_YERPE;	Q821S4, MUDD_CHLCV;	Q9PLG1, MUDD_CHLMU;
Q9Z701, MUDD_CHLPN;	O84767, MUDD_CHLTR;	

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